

174401	ATU036 341	ATL8C4 5437:1 ..834	nap					534:24 6..110	72				29	473) Contai ns simila rity to hypoth etical 43.1 KD protei n in NDK- GCPE interg enic region gb 493 519 from E. coli sequen ce gb U02 965. [Arabi dopsis thalia na]
								ATL8C4 5437:1 72...46 9	0.67	g30804 44	30	662	3.8e- 75	(AL022 605) putati ve protei n [Arabi dopsis thalia na]

174402	ATU036 342	ATL8C4 6738:1 125..1	nap				ATL8C4 6738:8 15..28 1	0.67	g37383 37	30	399	1.9e- 49	(AC005 170) putati ve revers e transc riptas e [Arabi dopsis thalia na]
174403	ATU036 343	ATL8C2 1773:1 ..1066	nap				ATL8C2 1773:8 9..826	0.67	g47739 08	30	291	1.8e- 19	(AF147 259) No defini tion line found [Arabi dopsis thalia na]
174404	ATU036 344	ATL8S4 460:1. .522	nap				ATL8S4 460:19 6..129	0.66	g41917 91	30	245	4.4e- 25	(AC005 917) putati ve sf21 {Helia nthus annuus } protei n [Arabi dopsis thalia na]
174405	ATU036	ATL8C6	nap				ATL8C6	0.66	g48039	30	332	9.7e-	(AC006

174411	ATU036 351	ATL8C5 009:59 4..1	nap						ATL8C5 009:51 1..205	0.63		g34829 19	30	537	1.4e- 33	e C [Nicot iana rustic al] (AC003 970) Putati ve protei n kinase [Arabi dopsis thalia na]
174412	ATU036 352	ATL8S2 8360:1 13..40 1	nap						ATL8S2 8360:1 22..39 5	0.63		g16522 28	30	185	3.1e- 15	(D9090 4) hypoth etical protei n [Synec hocyst is sp.]
174413	ATU036 353	ATL8C5 69:1.. 1428	nap						ATL8C5 69:13.. .1051	0.63		g39351 64	30	894	6.4e- 80	(AC004 557) F17L21 .7 [Arabi dopsis thalia na]
174414	ATU036 354	ATL8S1 7613:4 50..1	nap						ATL8S1 7613:4 05..92	0.62		g20784 83	30	95	0.999	(U4320 0) antifr eeze glycop eptide